

THE EFFECT OF MISSING DATA AND OF TWO SOURCES OF
CHARACTER VALUES ON A PHENETIC STUDY OF
THE WILLOWS OF CALIFORNIA

THEODORE J. CROVELLO

Numerical taxonomy is still in a phase of expansion and evaluation. New methods continue to appear and more and more taxonomists are using it to estimate phenetic relationships among plant taxa. Gilmartin (1967) provided an extensive bibliography of recent numerical taxonomic applications in botany. Sokal and Sneath (1963) review earlier works.

Concurrent with the development of numerical taxonomy, the use of computers in biological collections and floras has increased. Crovello (1967) lists the uses of electronic data processing in biological collections. Some uses do not involve measurements of characters on specimens, but for other purposes the measurements given in floras would be very helpful. Before information in floras can be employed, the reliability of the measurements provided in a floristic treatment and the problem of missing data must be investigated. By reliability of measurements I mean how accurate they are in estimating the parametric values (e.g., the mean and standard deviation) of a certain character in a given taxon. The problem of missing data arises because many floras are written without the prior preparation of a complete taxon by character table. Many floristic taxonomists include values of a character only in those taxa where the characters are diagnostic of it. This is brought about partly because preserved specimens rarely have all stages of the life cycle present.

The purpose of the present study is twofold: 1, to test the effect of missing data on a phenetic study of a group of plants, and 2, to ascertain the reliability of two sources of information about characters in taxospecies, one arising from a floristic treatment (Munz, 1959) and the other from a study of monographic proportions (Crovello, 1966). The latter is assumed to involve more measurements per character per taxospecies than the former. The results should be of value in estimating the reliability of information from floristic studies for estimation of phenetic relationships. This is especially timely in view of the proposed Flora North America Project. A natural byproduct will be further comprehension of the pattern of variation *among* the willows of California. By this I mean the phenetic relationships among (not *within*) the taxospecies in the context of the characters used. For example, Fig. 1 indicates relationships among the willows based on 43 characters. This figure is a reflection of the pattern of variation among the taxospecies based on these 43 characters.

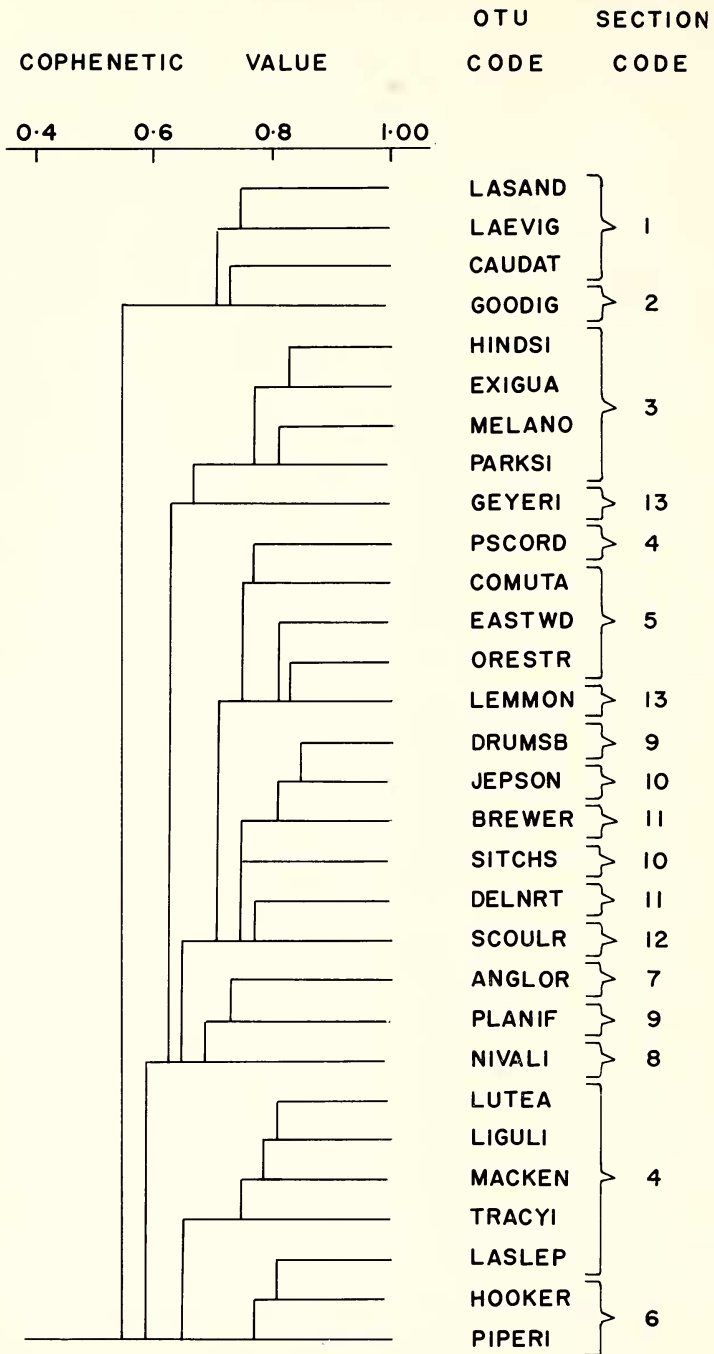


FIG. 1. Phenogram of analysis using data from Crovello (1966) for 43 characters.

MATERIALS AND METHODS

In his floristic treatment of *Salix*, Munz (1959) used 57 morphological characters, but eight were invariant (did not distinguish any willow species) and were omitted. Of the remaining 49 characters, Crovello (1966) could not obtain information on six of them, so both studies had only 43 characters in common.

Six analyses were made. These differed only in the input data that were used in each. They are the following: 1, data from Crovello (1966) on the 43 available characters of the 57 characters used by Munz (1959); 2, data from Crovello (1966) as the previous analysis, but with the same pattern of missing information as found in Munz (1959); 3, data from Munz (1959) on 43 characters that Crovello (1966) also scored; 4, data from Munz (1959) on 49 of the 57 characters used by him; 5, data from Crovello (1966) on 131 characters; 6, data from Crovello (1966) on 202 characters. Analyses 5 and 6 were included to serve as standards with which to compare the results of the first four analyses. Two standards are considered more desirable, since no one result that uses a large number of characters can be taken as depicting the true, overall, phenetic relationships better than another analysis using many characters. With two "standards" before him, the reader has some idea of the variability of results even when information is available on over 100 characters. Table 1 lists the characters employed in the first four analyses. Crovello (1966) gives the characters used in the last two analyses. In it he treated floral characters common to both sexes, e.g., ament length, as one character, but in the present study these are treated as two characters.

For analyses 3 and 4, the information contained in the treatment of the genus by Munz (1959) was used exclusively. This includes the information located both in the keys and in the description of each species. The information under each variety was incorporated into the description of its species. The other four analyses used information gathered by Crovello (1966). He used herbarium specimens to reinforce his personal collections. From 7 to 15 plants were chosen for each of the 31 taxospecies recognized by Munz (1959) as native to California. Most plants were represented by several herbarium sheets. Crovello (1966) provides a list of specimens used. This list and copies of the taxospecies by character tables, or Basic Data Matrices (BDM's) are on file on punched cards in The Herbarium, University of Notre Dame.

Crovello (1966) concluded that *Salix coulteri* is a synonym of *Salix sitchensis*. As a result, the number of taxospecies analyzed in the present study is 30, one less than the number recognized by Munz. Table 2 lists the 30 taxospecies and their codes. They are grouped into sections according to the ideas of Schneider (1921).

There exists no one method of numerical taxonomy. The present study used only one because we are interested in the effect of different sources of data and not in the effect of different taximetric methods. Sokal and Sneath (1963) discuss a number of different methods.

TABLE 1. CHARACTERS USED IN THE PRESENT STUDY.
THE FIRST 43 WERE USED BY CROVELLO.

1. Plant habit	30. Stigma lobe length
2. Plant height	31. Style length
3. Last year's twig color	32. Capsule length
4. This year's twig color	33. Capsule pedicel length
5. Stipules present or absent	34. Stamen number
6. Stipule length	35. Stamen filament divided
7. Stipule shape	36. Stamen filament pubescent
8. Stipule margin	37. Presence or absence of pubescence on last year's trig
9. Petiole length	38. Presence or absence of pubescence on this year's twig
10. Petiole glandular	39. Presence or absence of pubescence on abaxial leaf surface
11. Blade length	40. Presence or absence of pubescence on adaxial leaf surface
12. Blade shape	41. Presence or absence of pubescence on adaxial side of floral scale
13. Blade margin entire	42. Presence or absence of pubescence on adaxial side of floral scale
14. Blade margin glandular	43. Presence or absence of pubescence on capsule surface
15. Blade margin revolute	44. Bark texture
16. Blade base shape	45. Bark color
17. Blade apex shape	46. Blade color
18. Blade veins prominent below	47. Time of flowering compared to time of leaf break
19. Blade abaxial side glaucous	48. Ovary shape
20. Blade adaxial side lustrous	49. Flowering period
21. Blade width	
22. Female ament length	
23. Male ament length	
24. Female peduncle length	
25. Female peduncle leaf number	
26. Female ament dense or lax	
27. Female floral scale length	
28. Female floral scale shape	
29. Female floral scale color	

The procedure used here was the same for each of the six analyses. For each analysis the raw data appears in the form of a taxospecies by character table, or Basic Data Matrix (BDM). Each character was transformed by condensation to remove the effect of weighting due to measurement of different characters in different units. For example, leaf length was measured in millimeters and leaf base shape was measured in angles. To give each character equal weight, each value of a certain character in the tables was condensed, i.e., the value of a character in a certain taxospecies was replaced by a value X_{ej} ,

$$X_{ej} = \frac{X_j - X_{\min}}{X_{\max} - X_{\min}}$$

where X_{ej} is the condensed value of character X in taxospecies j , X_j is the original value of character X in taxospecies j and X_{\min} and X_{\max} are the minimum and maximum observed values of character X in the BDM.

Next, I calculated the similarity between each pair of Operational Taxonomic Units (OTU's), here the taxospecies. I used a modification of the distance coefficient introduced by Sokal (1961). Call this modifi-

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To JOHN THOMAS HOWELL, indefatigable collector and student of the plants of California and the Galápagos Islands, and long time member of the California Botanical Society, this volume of Madroño is dedicated.

Tom Howell was born in Merced where he lived until he graduated from high school. It was as a student at the University of California in Berkeley, where he came under the influence of Professor W. L. Jepson, that he developed his interest in botany. Following his university graduation and subsequent completion of a master's degree, he went to southern California, where he spent the first two years of his professional life. In 1929 at the invitation of Miss Alice Eastwood he came to the California Academy of Sciences in San Francisco where he has remained.

As a collector of botanical specimens, Tom Howell has few equals, and his collection numbers have surpassed 40,000. Moreover, his collections have never lain idle to collect dust but he has determined them and incorporated them in the Herbarium of the Academy. Although most of his collecting has been done in California and the Galápagos Islands he has made important collections in several western states. In addition to his own collections which have added so extensively to the Academy's herbarium he has encouraged many others, both amateur and professional, to make collections which also have come to the Academy.

His numerous botanical papers have contributed greatly to our knowledge of the plants of the areas where he has collected and particularly his floras of several regions in northern and central California have contributed to a greater botanical understanding of local areas.

Leaflets of Western Botany, which he and Miss Eastwood began in 1932 and which he edited and published until the end of 1966, provided a vehicle for significant contributions to the botany of the western United States.

His many years of field work, his keen observations, his great enthusiasm and unlimited energy, have brought to him an unsurpassed knowledge of California's native and weedy plants. And he is always a source of botanical information, willingly, thoughtfully and kindly given, to all who seek him out.

CONTENTS

	PAGE
Frontispiece: John Thomas Howell	
A revision of <i>Petalonyx</i> (Loasaceae) with a consideration of affinities in subfamily Gronovioideae, <i>William S. Davis</i> and <i>Henry J. Thompson</i>	1
MacNab cypress in Northern California: A geographical review, <i>J. R. Griffin</i> and <i>C. O. Stone</i>	19
A new species of primrose from Nevada, <i>Noel H. Holmgren</i>	27
Reviews	30, 58, 94, 220, 287
Natural hybridization between <i>Encelia</i> and <i>Geraea</i> (Compositae) and some related experimental investigations, <i>Donald W. Kyhos</i>	33
New records of Myxomycetes from California. II, <i>Donald T. Kowalski</i>	43
Notes and News	46, 62, 78, 91, 95, 136, 164, 192, 210, 222, 234, 266, 288, 300
Cypress rusts in California and Baja California, <i>Roger S. Peterson</i>	47
Two new eriogonums from Baja California, Mexico, <i>James L. Reveal</i> and <i>Craig A. Hanson</i>	55
Check list of California Pteridophytes, <i>Lawrence L. Kiefer</i> and <i>Barbara Joe</i>	65
A new Hesperaloe from Sonora, Mexico, <i>Howard Scott Gentry</i>	74
Studies of juniper rusts in the West, <i>Roger S. Peterson</i>	79
A new species of <i>Pterigoneurum</i> from California, <i>Howard Crum</i>	92
The scientific writings of Willis Linn Jepson, <i>Lawrence R. Heckard</i> , <i>John Thomas Howell</i> , and <i>Rimo Bacigalupi</i>	97
On the <i>Hordeum jubatum</i> - <i>H. brachyantherum</i> question, <i>Wm. W. Mitchell</i>	108
Notes on marine algae of Washington and southern British Columbia. II, <i>Richard E. Norris</i> and <i>John A. West</i>	111
<i>Gossypium trilobum</i> : An addendum, <i>Paul A. Fryxell</i> and <i>Clifford R. Parks</i>	117
Lee Bonar	124
A hybrid hawkweed from the Olympic Mountains of Washington, <i>Arthur R. Kruckeberg</i>	126
A survey of the vascular plants above timberline on Mt. Hood, Oregon, <i>Lurel Hall</i>	130
Documented chromosome numbers of plants	134
Variation in the diploid <i>Gossypium</i> species of Baja California, <i>Lyle L. Phillips</i> and <i>Duncan Clement</i>	137
California boletes. III. The genus <i>Suillus</i> , <i>Harry D. Thiers</i>	148
Chromosome variation in ecotypes of <i>Larrea divaricata</i> in the North American desert, <i>Tien Wei Yang</i> and <i>Charles H. Lowe</i>	161
Cytological and evolutionary studies in palustres violets, <i>Marja Sorsa</i>	165
A second species of <i>Ammobroma</i> (Lennoaceae), in Sinaloa, Mexico, <i>Robert L. Dressler</i> and <i>Job Kuijt</i>	179
Chromosome numbers in the tribe Pappophoreae (Gramineae), <i>John R. Reeder</i> and <i>D. N. Singh</i>	183
Some additions to the Flora of Texas. IV, <i>Donovan S. Correll</i>	187
New taxa, combinations, and chromosome numbers in <i>Arctostaphylos</i> (Ericaceae), <i>Philip V. Wells</i>	193
Chromosome counts in section <i>Simiolus</i> of the genus <i>Mimulus</i> (Scrophulariaceae). VII. New numbers for <i>M. guttatus</i> , <i>M. cupreus</i> , and <i>M. tilingii</i> , <i>R. K. Vickery, Jr.</i> , <i>K. W. Crook</i> , <i>D. W. Lindsay</i> , <i>M. M. Mia</i> , and <i>W. Tai</i>	211
A rust on <i>Anemopsis californica</i> , <i>Paul D. Keener</i> and <i>Jack E. Hampton</i>	218
The vegetational composition of the south slope of Mt. Pinos, California, <i>Richard J. Vogl</i> and <i>Brian C. Miller</i>	225
Taxonomy, variation, and chorology of three chromosome races of the <i>Calamagrostis canadensis</i> complex in Alaska, <i>Wm. W. Mitchell</i>	235
New records of Myxomycetes from California. III, <i>Donald T. Kowalski</i> and <i>Dwayne H. Curtis</i>	246

Chromosome counts in section Simiolus of the genus <i>Mimulus</i> (Scrophulariaceae). VIII. Chromosomal homologies of <i>M. glabratus</i> and its allied species and varieties, <i>M. M. Mia</i> and <i>Robert K. Vickery, Jr.</i>	250
Identity of the oak tree at Live Oak Tank, Joshua Tree National Monument, California, <i>John M. Tucker</i>	256
Seed ecology of <i>Sequoiadendron giganteum</i> , <i>N. Stark</i>	267
A cryptotaxonomic study of a natural hybrid between <i>Agropyron cristatum</i> and <i>A. subsecundum</i> , <i>W. S. Boyle</i> and <i>A. H. Holmgren</i>	277
The water hyacinth in California, <i>Jane H. Bock</i>	281
Notes on <i>Oryzopsis hendersonii</i> (Gramineae), <i>Richard Spellenberg</i>	283
New species of <i>Eriogonum</i> from Utah, <i>James L. Reveal</i>	289
The effect of missing data and of two sources of character values on a phenetic study of the willows of California, <i>Theodore J. Covelto</i>	301
Index	315

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I certify that the statements made by me above are correct and complete.

JOHN H. THOMAS, Editor

November 15, 1968

TABLE 2. THE 30 TAXOSPECIES OF SALIX IN CALIFORNIA RECOGNIZED BY CROVELLO (1966). ARRANGEMENT BY SECTIONS FOLLOWS SCHNEIDER (1921)¹.

Section Code	Taxospecies Symbol	Taxospecies
1. <i>Pentrandrae</i> Dumort.	LASAND	<i>S. lasiandra</i> Benth.
	CAUDAT	<i>S. caudata</i> (Nutt.) Hell.
	LAEVIG	<i>S. laevigata</i> Bebb.
2. <i>Nigrae</i> Loudon	GOODIG	<i>S. gooddingii</i> Ball
3. <i>Longifoliae</i> Anderss.	HINDSI	<i>S. hindsiana</i> Benth.
	EXIGUA	<i>S. exigua</i> Nutt.
	MELANO	<i>S. melanopsis</i> Nutt.
4. <i>Cordatae</i> Barr.	PARKSI	<i>S. parksiana</i> Ball
	LUTEA	<i>S. lutea</i> Nutt.
	LIGULI	<i>S. ligulifolia</i> (Ball) Ball
	MACKEN	<i>S. mackenziana</i> (Hook.) Barr.
	PSCORD	<i>S. pseudocordata</i> Anderss.
	LASLEP	<i>S. lasiolepis</i> Benth.
	TRACYI	<i>S. tracyi</i> Ball
5. <i>Andenophyllae</i> Schneid.	COMUTA	<i>S. commutata</i> Bebb
	EASTWD	<i>S. eastwoodiae</i> Ckll.
	ORESTR	<i>S. orestera</i> Schneid.
6. <i>Chrysanthae</i> Koch	PIPERI	<i>S. piperi</i> Bebb
	HOOKEA	<i>S. hookeriana</i> Barr.
7. <i>Ovalifoliae</i> Rydb.	ANGLOR	<i>S. anglorum</i> Cham. var. <i>antiplasta</i> Schneid.
8. <i>Reticulatae</i> Fries	NIVALI	<i>S. nivalis</i> Hook.
9. <i>Phylicifoliae</i> Dumort.	PLANIF	<i>S. planifolia</i> Pursh var. <i>monica</i> (Bebb) Schneid.
	DRUMSB	<i>S. drummondiana</i> Barr. var. <i>subcoerulea</i> (Piper) Ball
10. <i>Sitchenses</i> Bebb	SITCHS	<i>S. sitchensis</i> Sans.
	JEPSON	<i>S. jepsonii</i> Schneid.
11. <i>Brewerianae</i> Schneid.	BREWER	<i>S. breweri</i> Bebb
	DELNRT	<i>S. delnortensis</i> Schneid.
12. <i>Discolores</i> Barr.	SCOULR	<i>S. scouleriana</i> Barr.
13. <i>Fulvae</i> Barr.	LEMMON	<i>S. lemmonii</i> Bebb
	GEYERI	<i>S. geyeriana</i> Anders.

¹ The only exception to Schneider's assignments is *S. jepsonii*. He placed it in section *Phylicifoliae*.

cation the similarity coefficient, s_{jk} . Then,

$$s_{jk} = 1 - \left[\frac{\sum_{i=1}^{n_{jk}} (X_{ij} - X_{ik})^2}{n_{jk}} \right]^{1/2}$$

where X_{ij} and X_{ik} are the values of character i in OTU's j and k , respectively, and n_{jk} is the total number of characters used in the particular comparison. If there were no missing data, i.e., if the OTU by OTU relevance (Sokal and Sneath, 1963) were always 1.0, then n_{jk} would be the same for all combinations of pairs of OTU's.

TABLE 3. SUMMARY STATISTICS OF THE SIX ANALYSES.

Analysis number and description	Similarity coefficient		OTU by OTU relevance		Cophenetic correlation coefficient
	\bar{Y}	s	\bar{Y}	s	
1. Crovello 43	.593	.086	.972	.037	.746
2. Crovello 43 with Munz's missing data pattern	.578	.098	.482	.082	.678
3. Munz 43	.590	.085	.495	.079	.741
4. Munz 49	.587	.082	.483	.075	.719
5. Crovello 131	.617	.077	.908	.032	.858
6. Crovello 202	.615	.072	.829	.080	.871

Each set of similarity coefficients from one BDM forms a 30 by 30 OTU by OTU table, or Basic Similarity Matrix (BSM). Each cell in it indicates how similar two taxospecies are in the context of the characters used and based on the source of the data for the BDM being analyzed. The BSM was then used to group OTU's by the unweighted pair-group method (Sokal and Sneath, 1963). This results in a phenogram, a hierarchic presentation of phenetic relationships among the OTU's, *in the context of the characters analyzed*.

RESULTS

Figures 1 to 4 and Tables 3 and 4 present the results of the four analyses. Figures 5 and 6 are standards with which to compare the phenograms of the analyses using Munz's characters. The last two represent the maximum information on California willows available to the author at the present time. Figure 5 is based on the morphological characters studied by Crovello (1966) but without the six (or seven) unit pubescence characters scored for each organ. Figure 6 includes the 62 unit pubescence characters. *Two* standards were used: 1, to increase comprehension of the results of the present paper; and 2, to emphasize that in numerical taxonomy any one result is not the ultimate truth. For ease of presentation of results, we shall compare Figs. 1 through 6 with the latest nonnumerical monograph, which I summarize in Table 2.

Figure 1 gives the analysis of 43 characters using full data from Crovello (1966) which produced seven clusters. Beginning at the top, the first cluster contains four taxospecies that are the only representatives in California of the subgenus *Pleiandrae*. The next four OTP's are the representatives of section *Longifoliae* appearing in California. GEYERI then joins this cluster. The next cluster of five taxospecies includes members of sections *Cordatae*, *Adenophyllae* and *Fulvae*, while the subsequent cluster includes members of sections *Phylicifoliae*, *Sitchenses*, *Brewerianae* and *Discolores* in a mixed pattern. This is followed by a cluster consisting of three high-altitude willows, ANGLOR, PLANIF and NIVALI. The next to last cluster includes four members of section *Cordatae*, while the last cluster contains LASLEP, a polymorphic member of sec-

TABLE 4. CORRELATION BETWEEN ALL PAIRS OF THE SIX SIMILARITY MATRICES (LOWER LEFT), AND CORRELATION BETWEEN ALL PAIRS OF THE SIX PHENOGRAMS (UPPER RIGHT). IN ALL CASES $n = 435$.

Analysis number and description	Analysis number					
	1	2	3	4	5	6
1. Crovello 43	1.000	.852	.606	.643	.715	.577
2. Crovello 43 with Munz's missing data pattern	.887	1.000	.455	.437	.460	.455
3. Munz 43	.583	.450	1.000	.947	.658	.573
4. Munz 49	.604	.460	.974	1.000	.739	.653
5. Crovello 131	.835	.669	.655	.670	1.000	.889
6. Crovello 202	.733	.585	.595	.614	.889	1.000

tion *Cordatae*, and the two taxospecies of section *Chrysanthae* that appear in California.

Turning to Fig. 2, which is based on Crovello's data but uses the pattern of missing data present in the Munz data, seven clusters also are seen here. The first is similar to that of Fig. 1. The next two clusters of three and four taxospecies resemble two from Fig. 1, but TRACYI is out of place. The next cluster includes section *Longifoliae*, but here GEYERI has split the four members of that section. The next cluster contains six taxospecies. Except for ANGLOR, it is similar to a cluster in Fig. 1. The final two clusters also have their counterparts in Fig. 1, with the exception of ANGLOR mentioned above.

Figure 3 is based on Munz's data on the 43 characters that are comparable to Crovello's. The first cluster of four taxospecies is the same as in previous figures. But then HINDSI appears as deviant from all other OTU's. The other three members of section *Longifoliae* are far removed from it. The next two clusters, the first with six taxospecies, bring together quite different OTU's as suggested by conventional taxonomy and by Figs. 1 and 2. ANGLOR is a dwarf alpine form, whereas SCOULR is a polymorphic shrub or small tree more common at lower altitudes. By inspection of the rest of Fig. 3 the reader can ascertain the similarities and differences of it when compared to the previous figures.

Figure 4 is based on the 49 characters used by Munz. Seven more or less distinct clusters emerge. The four OTU's of section *Longifoliae* are closer together now than in Fig. 3, but as in Fig. 2, GEYERI splits them. Here EASTWD also is among this cluster. Note at the bottom of Fig. 4 that ANGLOR is still grouped with SCOULR.

Figure 5 consists of eight clusters, the first three of which agree exactly with Fig. 1. The remainder of Figs. 1 to 4 is in less agreement with Fig. 5 and Fig. 6. Note, however, that Figs. 5 and 6 are not identical.

Table 3 gives summary statistics for the six analyses. Columns 1 and 2 list the mean and standard deviation of each of the Basic Similarity